

z219c-a_-r	MRVSGVLRLLALIFAIVTTWMFIRSYMS-FSMKTIRLPRWLASP---- <td>45</td>	45
MMU72677_1	-----	
219_HUMAN	MRLAGPLRIVVLVSVGVTWIVVSILLGGPGSGFPRIQQLFTSPESSVTA	50
D87120_1_	MRVAGAAKLVVAVAVFLLTFYVISQVFE- IKMDASLG NLFARSALDTAAR	49
z219c-a_-r	EIQVKKYKCGLIKPCPANYFAFKICSGAANVVGPTMCFEDRMIMSPVKNN	95
MMU72677_1	-----	
219_HUMAN	APRARKYKCGLPQPCPEEHLAFRVVSGAANVIGPKICLEDKMLMSSVKDN	100
D87120_1_	STKPPRYKCGISKACPEKHFAFKMASGAANVVGPKICLEDNVLMSGVKNN	99
z219c-a_-r	VGRGLNIALVNGTTGAVLGQKAFDMYSGDVMHLVKFLKEIPGGALVLVAS	145
MMU72677_1	-----TGQVMKKDSFDMYSGDPQLLLLNFLTEIPDSTLVLVAS	37
219_HUMAN	VGRGLNIALVNGVSGELIEARAFDMWAGDVNDLLKFIRPLHEGTLVFVAS	150
D87120_1_	VGRGINVALANGKTGEVLDTKYFDMWGGDVAPFIEFLKAIQDGTIVLMGT	149
z219c-a_-r	YDDPGTKMNDESRLFSDLGSSYAKQLGFRDSWVFIGAKDLRGKSPFEQF	195
MMU72677_1	YDDPGTKMNDKIKTLFSNLGSSYAKQLGFRDSWVFIGAKDLKSKSPYEQF	87
219_HUMAN	YDDPATKMNEETRKLFSSELGSRNAKELAFRDSWVFIGAKGVQNKSPFEQH	200
D87120_1_	YDDGATKLNDEARRLIADLGSTSITNLGFRDNWVFCGGKGIKTKSPFEQH	199
z219c-a_-r	LKNSPDNTKYE GWPELLEMEGCMPPKPF--	223
MMU72677_1	LKNNPETNKYD GWPELLELEGVPRKVM--	115
219_HUMAN	VKNSKHSNKYEGCPEALEMEGCIPRRSTAS	230
D87120_1_	IKNNKDTNKYEGWPEVVEMEGCIPQKQD--	227

Figure